

	20	40	60	80	100		
Consensus	-----					EGKGDGLKEGSLXNDTGLEDET NXAND- N- LXNRQQM- GNKKKQA- GGR	45
1.1	-----					MD . TVT . LAMA . GVEK . K . . . G . DR . . RETNQSE . E . - RK .	40
1.4	-----					MD . TVT . LAMA . GVEK G . DR . . RETNQSE . E . - RK .	40
1.6	-----					MD . TVT . LAMA . GVEK G . DR . . REPNQSE . E . - RK .	40
1.7	-----					MD . TVT . LAMA . GVEK G . DR . . RETNQSE . E . - RK .	40
1.8	-----					MD . TVT . LAMA . GVEK G . DR . . RETNQSE . E . - RK .	40
1.9	-----					MD . TVT . LAMA . GVEK G . DR . . RETNQSE . E . - RK .	40
1.11	-----					MD . TVT . LAMA . GVEK G . DR . . RETNQSE . E . - RK .	40
2.2	-----					MEQTKDGLMANDPGLDA . TRKEYE . . . D . . . RMVK . . . G . E . G . RA . . QR . S . V . . .	55
2.4	-----					MEQTKDGLMANDPGLDA . TRKEYE . . . D . . . RMVK . . . G . E . G . RA . . QR . S . V . . .	55
2.5	-----					MEQTKDGLMANDPGLDA . TRKEYE . . . D . . . RMVK . . . G . E . G . RA . . QR . S . V . . .	55
2.7	-----					MEQTKDGLMANDPGLDA . TRKEYE . . . D . . . RMVK . . . G . E . G . RA . . QR . S . V . . .	55
2.8	-----					MEQTKDGLMANDPGLDA . TRKEYE . . . D . . . RMVK . . . G . E . G . RA . . QR . S . V . . .	55
2.10	-----					MEQTKDGLMANDPGLDA . TRKEYE . . . D . . . RMVK . . . G . E . G . RA . . QR . S . V . . .	55
2.11	-----					MEQTKDGLMANDPGLDA . TRKEYE . . . D . . . RMVK . . . G . E . G . RA . . QR . S . V . . .	55
3.1	-----					MMDSQSNSTD L FDEAHGEN I TGRSGERQK . NSTGLRG . RQ . E . N . . Q . G . R . MV . K . ERQE . Q . . E - MQQQR - S - . K	73
3.3	-----					MMDSQSNSTD L FDEAHGEN I TGRSGERQK . NSTGLRG . RQ . E . N . . Q . G . R . MV . K . ERQE . Q . . E - MQQQR - S - . K	73
3.4	-----					MMDSQSNSTD L FDDAHGEN I TGRSGERQK . NSTGLQG . RQ . E . N . . Q . G . R . MV . K . ERQE . Q . . E - MQQQR - S - . K	73
S3.1	MQRHERNLKQDWMTR I EWLKTTDWK I D N ELDNGKAKMHGETA L F FWDKHKHEKKN TGLAGERQKGN GTTSQ E . RR I E . NN . QVR . . . TP . K . EQWGKQ . . E - RQERRS - . EK					108	
4.1	-----					MDTTEKENGQD TNRKRNSEQGNATNMLERTRNEDAGENSNETWATVVSKKKEGKG . GQE . K . N . QSIASI . S . K . TEGQM . A . SQK . . . LNKL RYQ .	101
4.3	-----					MDTTEKENGQD TNGKRNREQGNATNMLERTRNEDAGENSNETWATVVSKKKEGKG . GQE . K . N . QSIASI . S . K . TEGQM . A . SQK . . . LNKL RYQ .	101
4.4	-----					MDTTEKENGQD TNGKRNREQGNATNMLERTRNEDAGENSNETWATVVSKKKEGKG . GQE . K . D . QSIASI . SVK . TEGQR . A . SQK . . . LNKL RSQ .	101
4.5	-----					MDTTEKENGQD TNGKRNREQGNATNMLERTRNEDAGENSNETWATVVSKKKEGKG . GQE . K . N . QSIASI . S . K . TEGQM . A . SQK . . . LNKL RYQ .	101
4.6	-----					MDTTEKENGQD TNGKRNREQGNATNMLERTRNEDAGENSNETWATVVSKKKEGKG . GQE . K . N . QSIASI . S . K . TEGQM . A . SQK . . . LNKL RYQ .	101
4.9	-----					MDTTEKENGQD TNGKRNREQGNATNMLERTRNEDAGENSNETWATVVSKKKEGKG . GQE . K . D . QSIASI . SVK . TEGQR . A . SQK . . . LNKL RSQ .	101
4.10	-----					MDTMEKENGQD TNGKRNREQGNATNMLERTQNE DARENSNETWATVVSKKKEGKG . . QE . K . D . QSTAI I . SVK . TEGQR . A . SQK . . . LNKI RYQ .	101
4.11	-----					MDTMEKENGQD TNGKRNREQGNATNMLERTQNE DARENSNETWATVVSKKKEGKG . . QE . K . D . QSTASI . SVK . TEGQR . A . SQK . . . LNKI RYQ .	101
4.12	-----					MDTTEKENGQD TNEERNGEQGNATNMFERTRNEDAGENSNETWATVVSKKKEGKGCGQE . K . D . QSNASI . S . K . IEGQR . A . SQK . . . LNKL RYQ .	101
4.14	-----					MDTTEKENGQD TNGKRNREQGNATNMLERTRNEDAGENSNETWATVVSKKKEGKG . GQE . K . N . QSIASI . S . K . TEEQM . A . SQK . . . LNKL RYQ .	101
4.15	-----					MDTTEKENGQD TNGKRNREQGNATNMLERTRNEDAEENSNETWATVVSKKKEGKG . GQE . K . N . QSIASI . S . K . TEGQM . A . SQK . . . LNKL RYQ .	101
5.3	-----					MESTKDKFTGLSGKTRGENNT EMDGEKQRGDDTGVREGKQMG . N . A QGG . . . M . M - ERQE . . . Q - - ERQHER - R . K	73
5.5	-----					MESTKDKFTGLSGKTRGESNT EMDGEKQRGDDTGVREGKQMG . N . A RGG . . . M . M - ERQE . . . Q - - ERQHER - R . K	73
5.6	-----					MESTKDKFTGLSGKTRGENNT EMDGEKQRGDDTGVREGKQMG . N . A QGG . . . M . M - ERQE . . . Q - - ERQHER - R . K	73
6.1	-----					MADGCTATNTDMAGETRLVKDNLGAGGTRKA I D IGLGG . T . TA . . N . . AGG . SK . . E - . G . DK . PRNGNKTE . SKFAE .	78
6.2	-----					MADGCTATNTDMAGETRLVKDNLGAGGTRKA I D IGLGG . T . TA . . N . . AGG . SK . . E - . G . DK . PRNGNKTE . SKFAE .	78
6.3	-----					MADGCTATNTDMAGETRLVKDNLGAGGTRKA I D IGLGG . T . TA . . N . . AGG . SK . . E - . G . DK . PRNGNKTE . SKFAE .	78
6.5	-----					MADGCTATNTDMAGETRLVKDNLGAGGTRKA I D IGLGG . T . TA . . N . . AGG . SK . . E - . G . DK . PRNGNKTE . SKFAE .	78
6.6	-----					MADGYTATNTDMAGETRLVKDNLGAGGTRKA I D IGLGG . T . TA . . N . . AGG . SK . . E - . G . DK . PRNGNKTE . SKFAE .	78
6.7	-----					MADGYTATNTDMAGETRLVKDNLGAGGTRKA I D IGLGG . T . TA . . N . . AGG . SK . . E - . G . DK . PRNGNKTE . SKFAE .	78
7.1	-----					MEKQRGE . I V SMDGDDGQ . R .	21
7.3	-----					MEKQRGE . I V SMDGDDGQ . R .	21
7.4	-----					MEKQRGE . I V SLDGDDGQ . R .	21
7.5	-----					MEKQRGE . I V SMDGDDGQ . R .	21
7.6	-----					MEKQRGE . I V SMDGDDGQ . R .	21

	120	140	160	180	200	220					
Consensus	KYLKEATV	IINVEDVEEVRAEDI	IKAVTEKCGXGKI	LALRPKQNKEYELTMESEEECEKLLDGLM	INGENCEVKKLQNRD	YVVSFLHLPVYLEDEE	ILEKLDGWGVCPI SKI 157				
1.1	ITVDIGQAT	...M.V.....RI.D	...V...H...V.L.R.	DADE.T.E.T.K.I	A.R.....M.....VA.KD	.D...H..... 152				
1.4	ITVDIGQAT	...M.V.....RI.D	...V...H...V.L.R.	DADE.T.E.T.K.I	A.R.....M.....VA.KD	.D...Y..... 152				
1.6	ITVEIGQAT	...M.V.....RI.D	...V...H...V.L.R.	DADE.T.E.T.K.I	A.R.....M.....VA.KD	.D.F.H..... 152				
1.7	ITVDIGQAT	...M.V.....RI.D	...V...H...V.L.R.	DADE.T.E.T.K.I	A.R.....M.....VA.KD	.D...H..... 152				
1.8	ITVDIGQAT	...M.V.....RI.D	...V...H...V.L.R.	DADE.T.E.T.K.I	A.R.....M.....VA.KD	.D...Y..... 152				
1.9	ITVDIGQAT	...M.V.....RI.D	...V...H...V.L.R.	DADE.T.E.T.K.I	A.R.....M.....VA.KD	.D...Y..... 152				
1.11	ITVDIGQAT	...M.V.....RI.D	...VV...H...V.LKR.	DADE.T.E.T.K.I	A.R.....M.....VA.KD	.D...H..... 152				
2.2Q.....IH.....R.GT.L.Q.T.KV...IH...F...M.....K.....T..... 167				
2.4K.....IH.....R.GT.L.Q.T.KV...IH...F...M.....K.....T..... 167				
2.5Q.....IH.....R.GT.L.Q.T.KV...IH...F...M.....K.....T..... 167				
2.7K.....IH.....R.GT.L.Q.T.KV...IH...F...M.....K.....T..... 167				
2.8K.....IH.....R.GT.L.Q.T.KV...IH...F...M.....K.....T..... 167				
2.10Q.....IH.....R.GT.L.Q.T.KV...IH...F...M.....K.....T..... 167				
2.11Q.....IH.....R.GT.L.Q.T.KV...IH...F...M.....K.....T..... 167				
3.1	.M.....L.DK.KVLE.NL.....IRDD.DL.N.T.D	.M...IK.T...M.....M.A.D	QL.D...E...F...T... 185			
3.3	.M.....L.DK.KVLE.NL.....IRDD.DL.N.T.D	.M...IK.T...M.....M.A.D	QL.D...E...F...T... 185			
3.4	.M.....L.MDK.KVLE.NL.....IRDD.DL.N.T.D	.M...IK.T...M.....M.A.D	QL.D...E...F...T... 185			
S3.1	.K.....L.L.DKIKVL.NI.....H.....IR.E.QI.F.T	.T.D...K.K...S.....M.A	QI.D...E...F...DF 220			
4.1	E.K.....LTMMIKNSTINNED.VI.LIGR.N.N.FK...I	IN.....E.....CATEKMN.S.IQQQ.VSNT...LP 213
4.3	E.K.....LTMMIKNSTINNED.VI.LIGR.N.N.FK...I	IN.....E.....CATEKMN.S.IQQQ.SNT...LP 213
4.4	E.K.....LTMM.RI.NATINNED.VI.LGR.N.N.FK...I	IN.....E.M...CATEKMN.S.IQQQ.SNT...LP 213
4.5	E.K.....LTMMIKNSTINNED.VI.LIGR.N.N.FK...I	IN.....E.....CATEKMN.S.IQQQ.SNT...LP 213
4.6	E.K.....LTMMIKNSTINNED.VI.LIGR.N.N.FK...I	IN.....E.....CATEKMN.S.IQQQ.SNT...LP 213
4.9	E.K.....LTMM.RI.NATINNED.VI.LGR.N.N.FK...I	IN.....E.M...CATEKMN.S.IQQQ.SNT...LP 213
4.10	E.K.....LTMM.RI.NSTINNED.VI.LGR.N.N.FK...I	IN.....E.M...CATEKMN.S.IQQQ.SNT...LP 213
4.11	E.K.....LTMM.RI.NSTINNED.VI.LGR.N.N.FK...I	IN.....E.M...CATEKMN.S.IQQQ.SNT...LP 213
4.12	E.K.....LTMM.RI.NATINNED.VI.LGR.N.N.FK...I	IN.....E.M...CATEKMN.S.IQD.QQ.SNT...LP 213
4.14	E.K.....LTMMIKNSTINNED.VI.LIGR.N.N.FK...V	IN.....E.....CATEKMN.S.IQQQ.SNT...LP 213
4.15	E.K.....LTMMIKNSTINNED.VI.LIGR.N.N.FK...I	IN.....E.....CATGKMN.S.IQQQ.SNT...LP 213
5.3L.DK.KVS.....LH.RI.RKDI.L.ET.DI.T.EM.....M.....SVS.L... 185	
5.5L.DK.KVS.....LH.RI.RKDI.L.ET.DI.T.EM.....M.....SVS.L... 185	
5.6L.DK.KVS.....LH.RI.RKDI.L.ET.DI.T.EM.....M.....SVS.L... 185	
6.1N.L.KV.IMQ.MR.GK.....ET.VS.H.....NSIV.E...E.LT... 190		
6.2N.L.KV.IMQ.MR.GK.....ET.VS.H.....NSIV.E...E.LT... 190		
6.3N.L.KV.IMQ.MR.GK.....ET.VS.H.....NSIV.E...E.LT... 190		
6.5N.L.KV.IMQ.MR.GK.....ET.VS.H.....NSIV.E...E.LT... 190		
6.6N.L.KV.IMQ.MR.GK.....ET.VS.H.....NSIV.E...E.LT... 190		
6.7N.L.KV.IMQ.MR.GK.....ET.VS.H.....NSIV.E...E.LT... 190		
7.1	VFQ.KL..LVEIVGEDKITMELLK.R.DV--VIGCY.NPNE.GKI...K.KNSRIMEIN.NEMN.T.IIG..TE...KAA.PV 131	
7.3	VFQ.KL..LVEIVGEDKITMELLK.R.DV--VIGCY.NPNE.GKI...K.KNSLIMEIN.NEMN.T.IIG..TE...KAA.PV 131	
7.4	VFQ.KL..LVEIVGEDKITMELLK.R.DV--VIGCY.NPNE.GKI...KVKNSRIMEIN.NEMN.T.IIG..TE...KAA.PV 131	
7.5	VFQ.KL..LVEIVGEDKITMELLK.R.DV--VIGCY.NPNE.GKI...K.KNSRIMEIN.NEMN.T.IIG..TE...KAA.PV 131	
7.6	VFQ.KL..LVEIVGEDKITMELLK.R.DV--VIGCY.NPNE.GKI...K.KNSRIMEIN.NEMN.T.IIG..TE...KAA.PV 131	

	240	260	280	300	320				
Consensus	KRRFYPGTDI	EDGTRFLKVRFPKEVASLPYSTXF	ETAEGPQYFRVMHSHQVKT	CRLCMSPEHXVKDCPNFKCYKCE	EERGHFARDCTTVKCEPKRFLNKC	ECWMEEEEEEG- 268			
1.1	N	V.T	RI	D.V	D.R	KF.A	D.NKV	G	V-
1.4	N	V.T	RI	D.V	D.R	KF.A	D.NKV	G	V-
1.6	N	V.T	RI	D.V	D.R	KF.A	D.NKV	G	V-
1.7	N	V.T	RI	D.V	D.R	KF.A	D.NKV	G	V-
1.8	N	V.T	RI	D.V	D.R	KF.A	D.NKV	G	V-
1.9	N	V.T	RI	D.V	D.R	KF.A	D.NKV	G	V-
1.11	N	V.T	RI	D.V	D.R	KF.A	D.NKV	G	V-
2.2			KM.T	ML	E	I	R	N	G.GG-
2.4			KM.T	ML	E	I	R	N	G.GG-
2.5		E	KM.T	ML	E	I	R	N	G.GG-
2.7			KM.T	ML	E	I	R	N	G.GG-
2.8			KM.T	ML	E	I	R	N	G.GG-
2.10			KM.T	ML	E	H	I	R	N
2.11			KM.T	ML	E	H	I	R	N
3.1	V	E	Y	R	I	RL	Q	D	M
3.3	V	E	Y	R	I	RL	Q	D	M
3.4	V	E	Y	R	I	RL	Q	D	M
S3.1	L	E	H	I	A	I	KL	M	L
4.1	R	KYH	MVA	V	K	K	NVG	M	E
4.3	R	KYH	MVA	V	K	K	NVG	M	E
4.4	R	KYH	MVA	VR	K	K	NVG	M	E
4.5	R	KYH	MVA	V	K	K	NVG	M	E
4.6	R	KYH	MVA	V	K	K	NVG	M	E
4.9	R	KYH	MVA	VR	K	K	NVG	M	E
4.10	R	KYH	MVA	VR	K	K	NVG	M	E
4.11	R	KYH	MVA	VR	K	K	NVG	M	E
4.12	R	KYH	MVA	VR	K	K	NVG	M	E
4.14	R	KYH	MVA	V	K	K	NVG	M	E
4.15	R	KYH	MVA	V	K	K	NVG	M	E
5.3	T	E	I	T	K	VM	D	R	F
5.5	T	E	I	T	K	VM	D	R	F
5.6	T	E	I	T	K	VM	D	R	F
6.1	C			RL	H	R	Q	D	LL
6.2	C			RL	H	R	Q	D	LL
6.3	C			RL	H	R	Q	D	LL
6.5	C			RL	H	R	Q	D	LL
6.6	C		L	RL	H	R	Q	D	LL
6.7	C			RL	H	R	Q	D	LL
7.1	MW	A		K	TDT	K	K	LT	TEH
7.3	MW	A		K	TDT	K	K	LT	TEH
7.4	ML	N	A	K	TDT	K	K	LT	TEH
7.5	MW	A		K	TDT	K	K	LT	TEH
7.6	MW	A		K	TDT	K	K	LT	TEH

	340	360	380	400	420	440	
Consensus	EI QVSRQVHEGNN-KE-EEDEE-AN-Q--EERTE-EE-KLMENEG-M---X-E-EG---ENR-E-----GEEQNEENX-I ELSGEE-RQD-X--TXQ-EI XAXXKVR LMD FE						350
1.1	.Q..G..MY..D.IQS-...DKTTT..EKRTES.S.KLQEN.TN.-IDRVT..Q..TTWTQMDMT--DSLKSVL.AAELSDSNNKDLN.GQQED.FWTQMDMT-DSFQKALD						366
1.4	.Q..G..MY..D.IQS-...DKTTT..EKRTES.S.KLQEN.TN.-IDRVT..Q..TTWTQMDMT--DSLKSVL.AAELSDSNNKDLN.GQQED.FWTQMDMT-DSFQKALD						366
1.6	.Q..G..MY..D.IQS-...DKTTT..EKRTES.S.KLQEN.TN.-IDRVT..Q..TTWTQMDMT--DSLKSVL.AAELSDSNNKDLN.GQQED.FWTQMDMT-DSFQKALD						366
1.7	.Q..G..MY..D.IQS-...DKTTT..EKRTES.S.KLQEN.TN.-IDRVT..Q..TTWTQMDMT--DSLKSVL.AAELSDSNNKDLN.GQQED.FWTQMDMT-DSFQKALD						366
1.8	.Q..G..MY..D.IQS-...DKTTT..EKRTES.S.KLQEN.TN.-IDRVT..Q..TTWTQMDMT--DSLKSVL.AAELSDSNNKDLN.GQQED.FWTQMDMT-DSFQKALD						366
1.9	.Q..G..MY..D.IQS-...DKTTT..EKRTES.S.KLQEN.TN.-IDRVT..Q..TTWTQMDMT--DSLKSVL.AAELSDSNNKDLN.GQQED.FWTQMDMT-DSFQKALD						366
1.11	.Q..G..MY..D.IQS-...DKTTT..EKRTES.S.KLQEN.TN.-IDRVT..Q..TTWTQMDMT--DSLKSVL.AAELSDSNNKDLN.GQQED.FWTQMDMT-DSFQKALD						366
2.2	.D-.D.....E...NR-.D-.....AI-.....RT..E.....N-N-N-...KK-.....NE.....E-VQ...QD-G...W--SQM-...SDSF.TF.STV.						349
2.4	.D-.D.....E...NR-.D-.....AI-.....RT..E.....N-N-N-...KK-.....NE.....E-VQ...QD-G...W--SQM-...SDSF.TF.STV.						349
2.5	.D-.D.....E...R...D-.....AI-.....RT..E.....N-N-N-...KK-.....NE...KE-VQ...QD-G...W--SQM-...SDSF.TF.STM.						350
2.7	.D-.D.....E...NR-.D-.....AI-.....RT..E.....N-N-N-...KK-.....NE.....E-VQ...QD-G...W--SQM-...SDSF.TF.STV.						349
2.8	.D-.D.....E...NR-.D-.....AI-.....RT..E.....N-N-N-...KK-.....NE.....E-VQ...QD-G...W--SQM-...SDSF.TF.STV.						349
2.10	.DR-D.....E...R...D-.....AI-.....RT..E.....N-N-N-...GKK-.....NE.....E-VQ...QD-G...W--SQM-...SDSF.TF.STV.						350
2.11	.DR-D.....E...R...D-.....AI-.....RT..E.....N-N-N-...KK-.....NE.....E-VQ...QD-G...W--SQM-...SDSF.TF.STV.						350
3.1	...G.M...S-EK-...S-...EEEQTT.QT--IQTTE...I---TLKE.NAKK.KEK.TETD-SQQIEH.EE-VAWTPMDITSSFKNVLDVI.KEDL-.EQSKGTD						390
3.3	...G.M...S-EK-...S-...EEEQTT.QT--IQTTE...I---TLKE.NAKK.KEK.TETD-SQQIEH.EE-VAWTPMDITSSFKNVLDVI.KEDL-.EQSKGTD						390
3.4	...G.M...S-EK-...S-...EEEQTT.QT--IQTTE...I---TLKE.NAKK.KEK.TETD-SQQIEH.EE-VAWTPMDITSSFKNVLDVI.KEDL-.EQSKGTD						390
S3.1	.N...G.M...R-EN-...S-...EVEKTTA.V--QQTITD...I---T-DE.VTKGKK.K.T-D-IQQTEQ.ED-LVWTPMDITPSFKSLDAT.NEVL-.EQTKET.						423
4.1	.T-.ASLENTATE-RK-...DK-TG-EENNTE-...D--RNEQEDKE-...ELG-...ANK.EG-...PNNRQFA.EQNDEDT...NN-...EK-.RTEDNEIRAI NEVRAMD						416
4.3	.T-.ASLENTATE-RK-...DK-TG-EENNTE-...D--RNEQEDKE-...ELG-...ANK.EG-...PNNRQFA.EQNDEDT...NN-...EK-.RTEDNEIRAI NEVRAMD						416
4.4	.T-.ASLENTATE-RK-...DK-TG-EENNTE-...D--RNEQEDKE-...ELG-...ANK.EG-...PNNRQFA.EQNDEDT...NN-...EK-.RTEDNEIRAI NEVRAMD						416
4.5	.T-.ASLENTATE-RK-...DK-TG-EENNTE-...D--RNEQEDKE-...ELG-...ANK.EG-...PNNRQFA.EQNDEDT...NN-...EK-.RTEDNEIRAI NEVRAMD						416
4.6	.T-.ASLENTATE-RK-...DK-TG-EENNTE-...D--RNEQEDKE-...ELG-...ANK.EG-...PNNRQFA.EQNDEDT...NN-...EK-.RTEDNEIRAI NEVRAMD						416
4.9	.T-.ASLENTATE-RK-...DK-TG-EENNTE-...D--RNEQEDKE-...ELG-...ANK.EG-...PNNRQFA.EQNDEDT...NN-...EK-.RTEDNEIRAI NEVRAMD						416
4.10	.T-.ASLENTATE-RK-...DK-TG-EENNTE-...D--RNEQEDKE-...ELG-...ANK.EG-...PNNRQFA.EQNDEDT...NN-...EK-.RTEDNEIRAI NEVRAMD						416
4.11	.T-.ASLENTATE-RK-...DK-TG-EENNTE-...D--RNEQEDKE-...ELG-...ANK.EG-...PNNRQFA.EQNDEDT...NN-...EK-.RTEDNEIRAI NEVRAMD						416
4.12	.T-.ASLENTATE-RK-...DK-TG-EENNTE-...D--RNEQEDKE-...ELG-...ANK.EG-...PNNRQFA.EQNDEDT...NN-...EK-.RTEDNEIRAI NEVRAMD						416
4.14	.T-.ASLENTATE-RK-...DK-TG-EENNTE-...D--RNEQEDKE-...ELG-...ANK.EG-...PNNRQFA.EQNDEDT...NN-...EK-.RTEDNEIRAI NEVRAMD						416
4.15	.T-.ASLENTATE-RK-...DK-TG-EENNTE-...D--RNEQEDKE-...ELG-...ANK.EG-...PNNRQFA.EQNDEDT...NN-...EK-.RTEDNEIRAI NEVRAMD						416
5.3	RN.IN...KV..VEQN.G..Q.GVL.PKDLLK.IQNG.GIQEDKK.HKRN-NEDDIFC.E.H.E--ER.KM...A.ERLI.EER-.TLFKE.EKR.EEDNENIEMETMD						404
5.5	RN.I...KV..VEQN.G..Q.GVL.PKDLLK.IQNG.GIQEDKK.HKRN-NEDDIFC.E.H.E--ER.KM...A.ERLI.EER-.TLFKE.EKR.EEDNENIEMETMD						404
5.6	RN.IN...KV..VEQN.G..Q.GVL.PKDLLK.IQNG.GIQEDKK.HKRN-NEDDIFC.E.H.E--ER.KM...A.ERLI.EER-.TLFKE.EKR.EEDNENIEMETMD						404
6.1	.K..D...Q...K-E-D--AKLEGD.EE.ATQQR.IEQ.KN.T.EV--VDFNED-...TK-...GGN-QMMEI.FSD-M.EN.Q.KIE-QNKDMDY-.ED.KVDNMDKCMK						398
6.2	.K..D...Q...K-E-D--AKLEGD.EE.ATQQR.IEQ.KN.T.EV--VDFNED-...TK-...GGN-QMMEI.FSD-M.EN.Q.KIE-QNKDMDY-.ED.KVDNMDKCMK						398
6.3	.K..D...Q...K-E-D--AKLEGD.EE.ATQQR.IEQ.KN.T.EV--VDFNED-...TK-...GGN-QMMEI.FSD-M.EN.Q.KIE-QNKDMDY-.ED.KVDNMDKCMK						398
6.5	.K..D...Q...K-E-D--AKLEGD.EE.ATQQR.IEQ.KN.T.EV--VDFNED-...TK-...GGN-QMMEI.FSD-M.EN.Q.KIE-QNKDMDY-.ED.KVDNMDKCMK						398
6.6	.K..D...Q...K-E-D--AKLEGD.EE.ATQQR.IEQ.KN.T.EV--VDFNED-...TK-...GGN-QMMEI.FSD-M.EN.Q.KIE-QNKDMDY-.ED.KVDNMDKCMK						398
6.7	.K..D...Q...K-E-D--AKLEGD.EE.ATQQR.IEQ.KN.T.EV--VDFNED-...TK-...GGN-QMMEI.FSD-M.EN.Q.KIE-QNKDMDY-.ED.KVDNMDKCMK						398
7.1	SLDLYES.EDAITG..KDE.DEQ-EE-DIMV.RRAGEEVQEP--D-D-SQDAM--GKFPQ--KQVVQEE..LL-F.GG.NGKKGQEK-AN-.DGKEH-.ENQRGRD						338
7.3	SLDLYES.EDAITG..KDE.DEQ-EE-DIMV.RRAGEEVQEP--D-D-SQDAM--GKFPQ--KQVVQEE..LL-F.GG.NGKKGQEK-AN-.DGKEH-.ENQRGRD						338
7.4	SLDLYES.E.AITG..KDE.DEQ-EE-DIMV.RRAGEEVQEP--D-D-SQDAM--GKFPQ--KQVVQEE..LL-F.GG.NGKKGQEK-AN-.DGKEH-.ENQRGRD						338
7.5	SLDLYES.E.AITG..KDE.DEQ-EE-DIMV.RRAGEEVQEP--D-D-SQDAM--GKFPQ--KQVVQEE..LL-F.GG.NGKKGQEK-AN-.DGKEH-.ENQRGRD						338
7.6	SLDLYES.E.AITG..KDE.DEQ-EE-DIMV.RRAGEEVQEP--D-D-SQDAM--GKFPQ--KQVVQEE..LL-F.GG.NGKKGQEK-AN-.DGKEH-.ENQRGRD						338

	460	480	500	520	
Consensus	RXNL	---E-KE-QXKEIXDLESRXMENXNKEMVGXR-QRRRXEKIKPNLETAKKK-LXRIKLSNRYEE	-----	411	
1.1	TEET	-----GQSID.QA...GHLQKDG...TQ.KSAK...SL.....R..L.KDEEIECANKYELLKGLEDMD	--	439	
1.4	TEET	-----GQSID.QA...GHLQKDG...TQ.KSAK...SL.....R..L.KDEEIECANKYELLKGLEDMD	--	439	
1.6	TEET	-----GQSID.KQA...GHLQKDG...TQ.KSAK...SL.....R..L.KDEEIECANKYELLKGLEDMD	--	439	
1.7	TEET	-----GQSID.QA...GHLQKDG...TQ.KSAK...SL.....R..L.KDEEIECANKYELLKGLEDMD	--	439	
1.8	TEET	-----GQSID.QA...GHLQKDG...TQ.KSAK...SL.....R..L.KDEEIECANKYELLKGLEDMD	--	439	
1.9	TEET	-----GQSID.QA...GHLQKDG...TQ.KSAK...SL.....R..L.KDEEIECANKYELLKGLEDMD	--	439	
1.11	TEET	-----GQSID.QA...EHLQKDG...TQ.KSAK...SL.....V..R..L.KDEEIECANKYELLKGLEDMD	--	439	
2.2	KEE	----R-.TG.AEGMAND.ETR...VD.NK.ERGFL...TF.V...DM...V.TKNRAKYVNRYEVLRLGLEEDEEK	425		
2.4	KEE	----R-.TG.AEGMAND.ETM...VD.NK.ERGFL...TF.V...DM...V.TKNRAKYVNRYEVLRLGLEEDEEK	425		
2.5	KEE	----R-.TG.AEGMAND.ETR...VD.NK.ERGFL...TF.V...DM...V.TKNRAKYVNRYEVLRLGLEEDEEK	426		
2.7	KEE	----R-.TG.AEGMAND.ETM...VD.NK.ERGFL...TF.V...DM...V.TKNRAKYVNRYEVLRLGLEEDEEK	425		
2.8	KEE	----R-.TG.AEGMAND.ETM...VD.NK.ERGFL...TF.V...DM...V.TKNRAKYVNRYEVLRLGLEEDEEK	425		
2.10	KEE	----R-.TG.AEGMAND.ETR...VD.NK.ERGFL...TF.V...DM...V.TKNRAKYVNRYEVLRLGLEEDEEK	426		
2.11	KEE	----R-.TG.AEGMAND.ETR...VD.NK.ERGFL...TF.V...DM...V.TKNRAKYVNRYEVLRLGLEEDEEK	426		
3.1	.ETGKHEE	-V.KDKL.KRQTRRSKVLKSLK.TARKKGLKKGQL.SHNRYDSLRLG-GEEED	-----	450	
3.3	.ETGKHEE	-V.KDKL.KRQTRRSKVLKSLK.TARKKGLKKGQL.SHNRYDSLRLG-GEEED	-----	450	
3.4	.ETGKHEE	-V.KDKL.KRQTRRSKVLKSLK.TARKKGLKKGQL.SHNRYDSLRLG-GEEED	-----	450	
S3.1	.ETEKHEV	-T.KDNF.KRQTRRSKALSKM.TTRKKVFKKGQL.SHNRY.FLRSL-GEEED	-----	482	
4.1	FERGNYGL	--EKKEKGKKIKDLKSRKSGLNIEMVLQR.KI.R.E..ERM.MK.FE-KGDARNICLSDSDSE	-----	485	
4.3	FERGNYGL	--EKKEKGKKIKDLKSRKSGLNIEMVLQR.KI.R.E..ERM.RK.FE-KGDARNICLSDSDSE	-----	485	
4.4	FERGNYGL	--EKKEKGKKIKDLKSRKSGLNIEMVLQR.KI.R.E..ERM.RK.FE-KGDARNICLSDSDSE	-----	485	
4.5	FERGNYGL	--EKKEKGKKIKDLKSRKSGLNIEMVLQR.KI.R.E..ERM.RK.FE-KGDARNICLSDSDSE	-----	485	
4.6	FERGNYGL	--EKKEKGKKIKDLKSRKSGLNIEMVLQR.KI.R.E..ERM.RK.FE-KGDARNICLSDSDSE	-----	485	
4.9	FERGNYGL	--EKKEKGKKIKDLKSRKSGLNIEMVLQR.KI.R.E..ERM.RK.FE-KGDARNICLSDSDSE	-----	485	
4.10	FERGNYGL	--EKKEKGKKIKDLKSRKSGLNIEMVLQR.KI.R.E..ERM.RK.FE-KGDARNICLSDSDSE	-----	485	
4.11	FERGNYGL	--EKKEKGKKIKDLKSRKSGLNIEMVLQR.KI.R.E..ERM.RK.FE-KGDARNICLSDSDSE	-----	485	
4.12	FERGNYGL	--EKKEKGKKIKDLKSRKSGLNIEMVLQR.KI.R.E..ERM.RK.FE-KGDARNICLSDSDSE	-----	485	
4.14	FERGNCGL	--EKKEKGKKIKDLKSRKSGLNIEMVLQR.KI.R.E..ERM.RK.FE-KGDARNICLSDSDSE	-----	485	
4.15	FERGNYGL	--EKKEKGKKIKDLKSRKSGLNIEMVLQR.KI.R.E..ERM.RK.FE-KGDARNICLSDSDSE	-----	485	
5.3	NSQMGDN	.V.TG.RN.EKSN.IGRE.EIKMDREER.TI...TL.....ST.R.-T-TRMK.MNRYEVLKEMEGDDE	-----	482	
5.5	NSQMGDN	.V.TG.RN.EKSN.IGRE.EIKMDREER.TI...TL.....ST.R.-FT-TRMK.MNRYEVLKEMEGDDE	-----	482	
5.6	NSQMGDN	.V.TG.RN.EKSN.IGRE.EIKMDREER.TI...TL.....ST.R.-T-TRMK.MNRYEVLKEMEGDDE	-----	482	
6.1	VNLRRRTLK	-VKPNLDNVRKKAKAN.F.VLRSLE.GEDE	-----	436	
6.2	VNLRRRTLK	-VKPNLDNVRKKAKAN.F.VLRSLE.GEDE	-----	436	
6.3	VNLRRRTLK	-VKPNLDNVRKKAKAN.F.VLRSLE.GEDE	-----	436	
6.5	VNLRRRTLK	-VKPNLDNVRKKAKAN.F.VLRSLE.GEDE	-----	436	
6.6	VNLRRRTLK	-VKPNLDNVRKKAKAN.F.VLRSLE.GEDE	-----	436	
6.7	VNLRRRTLK	-VKPNLDNVRKKAKAN.F.VLRSLE.GEDE	-----	436	
7.1	FMTR	-I-L.TPLI.LE.EM.MSAGNIKMLKRK.K.TEEETTKK	-----	379	
7.3	FMTR	-I-L.TPLI.LE.EM.MSAGNIKMLKRK.K.TEEETTKK	-----	379	
7.4	FMTR	-I-L.TPLI.LE.EM.MSAGNIKMLKRK.K.TEEETTKK	-----	379	
7.5	FMTR	-I-L.TPLI.LE.EM.MSAGNIKMLKRK.K.TEEETTKK	-----	379	
7.6	FMTR	-I-L.TPLI.LE.EM..SAGNIKMLKRK.K.TEEETTKK	-----	379	

	20	40	60	80	100																																										
Consensus	-----MMFRFLI	SFLFLLMVLN	CVSFNARGL	MDKXKFEK	VKEKCKRKD	VICLQETNWKE	EEVMNDFK	KKXWQGD	ILYNNGDGK	SGRGVA	ILIKKDV	LNFKK	I	VYNDKEG	KCLX	106																															
1.1	---MFFMNV	MSFMNV	FIFF	GIL	IRM	ELR	C	KEIR	R	S	EMRLL	ENSGVLC	TIMI	110																										
1.4	MMFFMNV	MSFMNV	FIFF	GIFL	IRM	ELR	C	IEIR	R	S	EMRLL	ENSGVLC	TIMI	111																										
1.5	MMIFMNV	MSFMNV	FMFF	GIL	IRM	ELR	C	IEIR	R	S	EMRLL	ENSGVLC	TIMI	111																										
1.6	---MFFMNV	MSFMNV	FIFF	GIL	IRM	ELR	C	IEIR	R	S	EMRLL	ENSGVLC	TIMI	110																										
1.8	MMFFMNV	MSFMNV	FIFF	GIFL	IRM	ELR	C	IEIR	R	S	EMRLL	ENSGVLC	TIMI	111																										
1.9	MMFFMNV	MSFMNV	FIFF	GIFL	IRM	ELR	C	IEIR	R	S	EMRLL	ENSGVLC	TIMI	111																										
1.10	MMIFMNV	MSFMNV	FMFF	GIL	IRM	ELR	C	IEIR	R	S	EMRLL	ENSGVLC	TIMI	111																										
1.11	MMFFMNV	MSFMNV	FIFF	GIL	IRM	ELR	C	IEIR	R	S	EMRLL	ENSGVLC	TIMI	111																										
2.1	---MR	CIG	DIRN	EGH	ALIFRRRNYV	107																										
2.2	---MR	CIG	DIRH	EGH	ALIFRRRNYV	107																										
2.3	---MR	CIG	DIRH	EGH	ALFIFRRRNYV	107																									
2.5	---MR	CIG	DIRH	EGH	ALIFRRRNYV	107																										
2.7	---MR	CIG	DIRH	EGH	ALIFRRRNYV	107																										
2.8	---MR	CIG	DIRH	EGH	ALIFRRRNYV	107																										
2.10	---MR	CIG	DIRN	EGH	ALIFRRRNYV	107																										
2.11	---MR	CIG	DIRN	EGH	ALIFRRRNYV	107																										
S2.1	---MR	CIG	DIRN	EGH	ALIFRRRNYV	107																										
S2.2	---MR	CIG	DIRN	EGH	ALIFRRRNYV	107																										
3.1	-----MTVFFLLR	KNEEIINDIYEDFYSTANAQMSVIKQMI	104																										
3.3	-----MTVFFLLR	KNEEIINDIYEDFYSTANAQMSVIKQMI	104																										
3.4	-----MTVFFLLR	NNEEIINDIYEDFYSTANAQMSVIKQMI	104																										
4.5	-----MSNVCVLVSKCERLTCLTKCILKENIKEMRLNEFSNIEKNRGIFEKVELDKTNRIII	93																								
4.10	-----MSNVCVLVSKCERLTCLTKCILKENIKEMRLNEFSNIEKNRGIFEKVELDKTNRIII	93																								
4.11	-----MSNVCVLVSKCERLTCLTKCILKENIKEMRLNEFSNIEKNRGIFEKVELDKTNRIII	93																								
4.14	-----MSNVCVLVSKCERLTCLTKCMLKENIKEMRLNEFSNIEKNRGIFEKVELDKTNRIII	93																								
5.3	-----ILKMEGIRNMVDYEKEFLFILKNNV	106																								
5.4	-----ILSKMEGIRFNMVDYEKEFLFILKNNV	106																							
5.6	-----ILKMEGIRNMVDYEKEFLFILKNNV	106																								
5.7	-----ILSKMEGIRNMVDYEKEFLFILKNNV	106																								
5.8	-----ILKMEGIRNMVDYEKEFLFILKNNV	106																								
6.1	---MYFKYL	FLFLSFTLNVGDLKQNLNDGERRRQFRMVEIKDVEELA	107																					
6.2	---MYFKYL	FLFLSFTLNVGDLKQNLNDGERRRQFRMVEIKDVEELA	107																					
6.4	---MYFKYL	FLFLSFTLNVGDLKQNLNDGERRRQFRMVEIKDVEELA	107																					
6.5	---MYFKYL	FLFLSFTLNVGDLKQNLNDGERRRQFRMVEIKDVEELA	107																					
6.6	---MYFKYL	FLFLSFTLNVGDLKQNLNDGERRRQFRMVEIKDVEELA	107																					
6.7	---MYFKYL	FLFLSFTLNVGDLKQNLNDGERRRQFRMVEIKDVEELA	107																					
7.5	-----WVLL	FFILY	AMCDLKIACNREMERIRFTHSNDKSIVGEYHSQRACLGGIHNVQIHGNRLG	104

	120	140	160	180	200	220	
Consensus	I E V N Y E G K E M I I A N I H A P T E E K D K K E F N V L N K V L K X X K E I I M M G D F N T V F X K Q D M A D G M V F K S D T G R K E L K E L M K E N N L I D I W R E R N E E K R E F S R R Q I V G N F I C Q T R I D F						217
1.1	C . M E . . K . K V . M V . V N K . . . Y . . . R D Y . . H E R V . I S . L E . E T . K I . E . M . . . V Q T K . Y C						221
1.4	C . M E . . K . K V . M V . V N K . . . Y . . . R D Y . . H E R V . I S . L E . E T . K I . E . M . . . V Q T K . Y C						222
1.5	C . M E . . K . K V . M V . V N K . . . Y . . . R D Y . . H E R V . I S . L E . E T . K Q . I I . E . M . . . V Q T K . Y C						222
1.6	C . M E . . K . K V . M V . V N K . . . Y . . . R D Y . . H E R V . I S . L E . E T . K I . E . M . . . V Q T K . Y C						221
1.8	C . M E . . K . K V . M V . V N K . . . Y . . . R D Y . . H E R V . I S . L E . E T . K I . E . M . . . V Q T K . Y C						222
1.9	C . M E . . K . K V . M V . V N K . . . Y . . . R D Y . . H E R V . I S . L E . E T . K I . E . M . . . V Q T K . Y C						222
1.10	C . M E . . K . K V . M V . V N K . . . Y . . . R D Y . . H E R V . I S . L E . E T . K Q . I I . E . M . . . V Q T K . Y C						222
1.11	C . M E . . K . K V . M V . V N K . . . Y . . . R D Y . . H E R V . I S . L E . E T . K I . E . M . . . V Q T K . Y C						222
2.1 H . L D Y . E G F R I Q . N I S R						218
2.2 H . L D Y . E G F R I N I S R						218
2.3 H . L D Y . E G F R I N I S R						218
2.5 H . L D Y . E G F R I E N I S . V R V						218
2.7 H . L D Y . E G F R I N I S R						218
2.8 H . L D Y . E G F R I N I S R						218
2.10 H . L D Y . E G F R I Q . N I S . V R						218
2.11 H . L D Y . E G F R I Q . N I S R						218
S2.1 H . L D Y . E G F R I Q . N I S R						218
S2.2 H . L D Y . E G F R I Q . N I S R						218
3.1	L . I . K D I . L V . V N . S E . S I R R F . N Y . Q . V C S R . I T E S . I E . K G M G K . K V N . S Y						215
3.3	L . I . K D I . L V . V N . S E . S I R R F . N Y . Q . V C S R . I T E S . I E . K G M G K . K P V N . S Y						215
3.4	L . I . K D I . L V . V N . S E . S I R R F . N Y . Q . V C S R . I T E S . I E . K G M G K . K V N . S Y						215
4.5	V K . L . G S F R V C N Y N . Y K E I D . L I E . N E N . F I L I A M Q R I . I D . S . N . R T . R N . Q N . R K C D . V . V G M Y E . A N R I L K . S L						204
4.10	V K . L . G S F R V C N Y N . Y K E I D . L I E . N E N . F I L I A M Q R I . I D . S . N . R T . R N . Q N . R K C D . V . V G M Y E . A N R I L K . S L						204
4.11	V K . L . G S F R V C N Y N . Y K E I D . L I E . N E N . F I L I A M Q R I . I D . S . N . R T . R N . Q N . R K C D . V . V G M Y E . A N R I L K . S L						204
4.14	V K . L . G S F R V C N Y N . Y K E I D . L I E . N E N . F I L I A M Q R I . I D . S . N . R T . R N . Q N . R K C D . V . V G M Y E . A N R I L K . S L						204
5.3 Q K I I E R Y . K V N A I N . K D						217
5.4 Q K I I E R H . K V N A I N . K D						217
5.6 Q K I I E R Y . K V N A I K . K D						217
5.7 Q K I I E R H . K V N A I N . K D						217
5.8 Q K I I E R Y . K V N A I N . K D						217
6.1	R M R - . . N N . T L C V A . . . E . D R I . D L M I L W G . T . L I N R I . L R N Q K . . A . H K V V V G Y L . D G N L K . S						216
6.2	R M R - . . N N . T L C V A . . . E . D R I . D L M I L W G . T . L I N R I . L R N Q K . . A . H K V V V G Y L . D G N L K . S A						216
6.4	R M R - . . N N . T L C V A . . . E . D R I . D L M I L W G . T . L I N R I . L R N Q K . . A . H K V V V G Y L . D G N L K . S						216
6.5	R M R - . . N N . T L C V A . . . E . D R I . D L M I L W G . T . L I N R I . L R N Q K . . A . H K V V V G Y L . D G N L K . S						216
6.6	R M R - . . N N . T L C V A . . . E . D R I . D L M I L W G . T . L I N R I . L R N Q K . . A . H K V G V G Y L . D G N L K . S						216
6.7	R M R - . . N N . T L C V A . . . E . D R I . D L M I L W G . T . L I N R I . L R N Q K . . A . H K V G V G Y L . D G N L K . S						216
7.5	. D F M . Y N E S F R L L . L Y . . N I . T E R . . V . K T M K P L C T G - - N C . V V V W C T R L . V S K S I N A S . R Y . N Q S E . M V . A E . P F V G V L K . S L						213

	240	260	280	300	320	
Consensus	I L C T R N I E N F I E N I Q Y E E T S L S D H K L L H F K V N X E N I Q K G P G X W I L N T T I L K N E D Y V Q K V K E I I E K E K E N E M Y D E D K R I W W E N T K Y Q I K K Y T I K Y C A V L Q R C K K Y K E K E V K K					328
1.1	V G N K K P F M L D W S V K R V V A V L S Q G I N M V L V F R Q N L E					332
1.4	V G N K K P F M L D W S V K R V V A V L S Q G I N M V L V F R Q N L E					333
1.5	V G N K K P F M L D W S V K R V V A V L S K Q G I N M V L V F R Q N L E					333
1.6	V G N K K P F M L D W S V K R V V A V L S Q G I N M V L V F R Q N L E					332
1.8	V G N K K P F M L D W S V K R V V A V L S Q G I N M V L V F R Q N L E					333
1.9	V G N K K P F M L D W S V K R V V A V L S Q G I N M V L V F R Q N L E					333
1.10	V G N K K P F M L D W S V K R V V A V L S K Q G I N M V L V F R Q N L E					333
1.11	V G N K K K P F M L D W S V K R V V A V L S Q G I N M V L V F R Q N L E					333
2.1	N F I T Q N R R T R					329
2.2	N F I T Q N R R T R					329
2.3	N F I T Q N R R T R					329
2.5	N F I T Q N R R T R					329
2.7	N F I T Q N R R T R					329
2.8	N F I T Q N R R T R					329
2.10	N F I T Q N R R T R					329
2.11	N F I T Q N R R T R					329
S2.1	N F I T Q N R R T R					329
S2.2	N F I T Q N R R T R					329
3.1	V I Y K R D V F V F N F T E R V T S E K G Q I N L V F L S L T F R I I K N R H T I E					326
3.3	V I Y K S D V F V F N F T E R V T S E K G Q I N L V F L S L T F R I I K N R H T I E					326
3.4	V I Y K R D V F V F N F T D E R V T S D N K I T V Q I N L V F L S L T F R I I K N R H T I E					326
4.5	F K K K E V Q Y V N T L K T Y E D F W I T M F N E E V E L I S K M D I E N I N S V N D F E T E T G L D L N K E S N I S E K I K A F N K I R					315
4.10	F K K E V Q Y V N T L K T Y E D F W I T M F N E E V E L I S K M N I E N I N S V N D F E T E T G L D L N R E S N I S E K I K A F N K I R					315
4.11	F K K K E V Q Y V N T L K T Y E D F W I T M F N E E V E L I S K M N I E N I N S V N D F E T E T G L D L N R E S N I S E K I K A F N K I R					315
4.14	F K K K E V Q Y V N T L K T Y E D F W I T M F N E E V E L I S K M D I E N I N S V N D F E T E T G L D L N R E S N I S E K I K A F N K I R					315
5.3	N F V D T R I R T N					328
5.4	N F V D T R I R T N					328
5.6	N F V D T R I R T N					328
5.7	N F V D T R I R T N					328
5.8	N F V D T R I R T N					328
6.1	K E K L G D Q S V Y F K T F L M I D F N E T K R K I E G P R K A I V D L N T I Q H D E M S K S I K A A H V R					327
6.2	K E K L G D Q S V Y F K T F L M I D F N E T K R K I E G P R K A I V D L N T I Q H D E M S K S I K A A H V R					327
6.4	K E K L G D Q S V Y F K T F L M I D F N E T K R K I E G P R K A I V D L N T I Q H D E M S K S I K A A H V R					327
6.5	K E K L G D Q S V Y F K T F L M I D F N E T K R K I E G P R K A I V D L N T I Q H D E M S K S I K A A H V R					327
6.6	K E K L G D Q S V Y F K T F L M I D F N E T K R K I E G P R K A I V D L N T I Q H D E M S K S I K A A H V R					327
6.7	K E K L G D Q S V Y F K T F L M I D F N E T K R K I E G P R K A I V D L N T I Q H D E M S K S I K A A H V R					327
7.5	C V K D L Q Y V K V S K F I E I G A A M I V M S L N L E S R G M C A S L E A R E S I E R C R Y M P L F E N N V C E D L E K I R S R S K Q R N F K R M E M E S M R N					324

	340	360	380	400	420	440			
Consensus	SLEKELNKEN	-- KDI EK I KELEQKLRD	LEEDKYKGAMLR	SRAKYTV	EGEKCTKFFFD	LEKQRGKAEI	I KEI KGRNGKIVKGNXEI	LEEI XHFYEDLFKAKGIDEEKERKLL	437
1.1	K..N..KN..	-G.N.Q....	QGR.NEM..	E.FE..R..K..F.....	R...S.M.R..RSK..N..EQHE.....	RSY..K..CTE..K..KEKGE..			442
1.4	K..N..KN..	-G.N.Q....	QGR.NEM..	E.FE..R..K..F.....	R...S.M.R..RSK..N..EQHE.....	RSY..K..CTE..K..KEKGE..			443
1.5	K..T..KN..	-G.N.Q....	QGR.NEM..	E.FE..R..K..F.....	R...S.M.R..RSK..N..EQHE.....	RSY..K..CTE..K..KEKGE..			443
1.6	K..N..KN..	-G.N.Q....	QGR.NEM..	E.FE..R..K..F.....	R...S.M.R..RSK..N..EQHE.....	RSY..K..CTE..K..KEKGE..			442
1.8	K..N..KN..	-G.N.Q....	QGR.NEM..	E.FE..R..K..F.....	R...S.M.R..RSK..N..EQHE.....	RSY..K..CTE..K..KEKGE..			443
1.9	K..N..KN..	-G.N.Q....	QGR.NEM..	E.FE..R..K..F.....	R...S.M.R..RSK..N..EQHE.....	RSY..K..CTE..K..KEKGE..			443
1.10	K..T..KN..	-G.N.Q....	QGR.NEM..	E.FE..R..K..F.....	R...S.M.R..RSK..N..EQHE.....	RSY..K..CTE..K..KEKGE..			443
1.11	K..N..KN..	NG.N.Q....	QGR.NEM..	E.FE..R..K..F.....	R...S.M.R..RSK..N..EQHE.....	RSY..K..CTE..K..KEKGE..			444
2.1							G.L.....A...I.....K.....I.....		438
2.2							S.....G.L.....A...I.....KY.....I.....		438
2.3							S.....G.L.....A...I.....K.....I.....		438
2.5							S.....T.....G.L.....A...I.....K.....E.....I.....		438
2.7							S.....G.L.....MA...I.....K.....NQ.....I.....		438
2.8							S.....G.L.....MA...I.....K.....NQ.....I.....		438
2.10							S.....G.L.....A...I.....K.....I.....		438
2.11							S.....G.L.....A...I.....K.....I.....		438
S2.1							S.....G.L.....A...I.....K.....I.....		438
S2.2							S.....G.L.....A...I.....K.....I.....		438
3.1	N..I..D.NE-	-Q....M.G..KEI..K..E..R..K..V.....	R.KA...T..I.Q.AK.ESIE..E..K..KKY..E..TQ.V..VQMS..						435
3.3	N..I..E.NE-	-Q....M.G..KEI..K..E..R..K..V.....	R.KA...T..I.Q.AK.ESIE..E..K..KKY..E..TQ.V..VQMS..						435
3.4	N..I..G.NE-	-Q....M.G..KEI..K..E..R..K..VA.....	R.KA...T..I.Q.AK.ESIE..E..K..KKY..E..TQ.V..VQMA..						435
4.5	EWNE.MS.V.-	GINVD..I..QE..KKI..E.C..VII..DI...RS..Y.YE..T.QR.A..C..TQD..T.EDKEG..N.TRR..KE..TEN.VNLND.HF..							425
4.10	EWNE.MS.V.-	GINVD..I..QE..KKI..E.C..VII..DI...RS..Y.YE..T.QR.AV..C..TQD..T.EDKEG..N.TRR..KE..TEN.VNLND.HF..							425
4.11	EWNE.MS.V.-	GINVD..I..QE..KKI..E.C..VII..DI...RS..Y.YE..T.QR.A..C..TQD..T.EDKEG..N.TRR..KE..TEN.VNLND.HF..							425
4.14	EWNE.MS.V.-	GINVD..I..QE..KKI..E.C..VII..DI...RS..Y.YE..T.QR.A..C..TQD..T.EDKEG..N.TRR..KE..TEN.VNIND.HF..							425
5.3							S.....G.L.....A...I.....K.....E.....I.....		437
5.4							S.....G.L.....A...I.....K.....E.....I.....		437
5.6							S.....G.L.....A...I.....K.....E.....I.....		437
5.7							S.....G.L.....A...I.....K.....E.....I.....		437
5.8							S.....G.L.....A...I.....K.....E.....I.....		437
6.1	E.NDS..Q.K.-V..R.LF..EE..NI..R.CR..I.....CI.D.....N..SK.NG..R.L.KE..EKI.ETQD..K.VRD..S...KGE.N.KE.KN..								436
6.2	E.NDS..Q.K.-V..R.LF..EE..NI..R.CR..I.....I.D.....N..SK.NG..R.L.KE..EKI.ETQD..K.VRD..S...KGE.N.KE.KN..								436
6.4	E.NDS..Q.K.-V..R.LF..EE..NI..R.CR..I.....I.D.....N..SK.NG..R.L.KE..EKI.ETQD..K.VRD..S...KGE.N.KE.KN..								436
6.5	E.NDS..Q.K.-V..R.LF..EE..NI..R.CR..I.....I.D.....N..SK.NG..R.L.KE..EKI.ETQD..K.VRD..S...KGE.N.KE.KN..								436
6.6	E.NDS..Q.K.-V..R.LF..EE..NI..R.CR..I.....I.D.....N..SK.NG..R.L.KE..EKI.ETQD..K.VRD..S...KGE.N.KE.KN..								436
6.7	E.NDS..Q.K.-V..R.LF..EE..NI..R.CR..I.....I.D.....N..SK.NG..R.L.KE..EKI.ETQD..K.VRD..S...KGE.N.KE.KN..								436
7.5	N...I.E.NPE--	YTK.NFLRIKGV.EMY.TE.CE..KV...T..AL...R...Y.LN...GKQRKNY.T.L.NE..EKITDYVK...TVQL...R...KENTKQ.YVD.V..							433

	460	480	500	520	540	
Consensus	NYIKVKLEKDDKKECDREIXEEIEIAINQLNKKKSPGIDGLGSEFYIVFKDILKXILKEVFKEIFKCKEMNERMGMLMGLIYKRKGAKTELXNYRPI TMLNTDLKILAK					548
1.1	.L.SRV.EGE.R.E.R.KR.SG.....YDD.ENG.I.K.....V.G.D.VA.K.....					553
1.4	.L.SRV.EGE.R.E.R.KR.SG.....SS.G.YD.ENG.IIK.....V.G.D.VD.K.....					554
1.5	.L.SRV.EGE.R.E.R.KR.SG.....SS.YD.ENG.I.K.....V.G.D.VA.K.....					554
1.6	.L.SRV.EGE.R.E.R.KR.SG.....Y.SS.YD.ENG.I.K.....V.G.D.VD.K.....					553
1.8	.L.SRV.EGE.R.E.R.KR.SG.....SS.G.YD.ENG.IIK.....V.G.D.VD.K.....					554
1.9	.L.SRV.EGE.R.E.R.KR.SG.....SS.G.YD.ENG.IIK.....V.G.D.VD.K.....					554
1.10	.L.SRV.EGE.R.E.R.KR.SG.....SS.YD.ENG.I.K.....V.G.D.VA.K.....					554
1.11	.L.SRV.EGE.R.E.R.KR.SG.....SS.YD.ENG.I.K.....V.G.D.VA.K.....					555
2.1H.N.....E.....I.N.....G.....D.....Q.....					549
2.2I.N.....E.....K.....I.N.....G.....Q.....					549
2.3N.....D.D.T.....I.N.....G.....Q.....					549
2.5N.....E.....K.....I.N.....G.....T.....Q.....					549
2.7N.....D.....I.N.....G.....Q.....					549
2.8N.....D.....I.N.....G.....Q.....					549
2.10N.....E.....I.N.....G.....D.....Q.....					549
2.11N.....E.....I.N.....G.....D.....Q.....					549
S2.1N.....E.....I.N.....G.....D.....Q.....					549
S2.2N.....E.....I.N.....G.....D.....Q.....					549
3.1	KQ.T.IDEE...Q.G.RK.ES.....N.VC.E.IP.TD...LQKE.L.....H...E.L.K.....T					545
3.3	KQ.T.IDEE...Q.G.RK.ES.....N.VC.E.IP.TD...LQKE.L.....H...E.L.K.....T					545
3.4	KQ.T.VDEE...Q.G.RK.ES.....N.VC.E.IP.ID...LQKE.L.....H...E.L.K.....T					545
4.5	EK.TA.N.E.EM.ES.TDL.T.D.SRNG.....PV.K.NV.IP.N.IYSD.YEKG.LTNS.KK.MV.M.....N.D.GD.K...LS...Y...					536
4.10	EK.TA.N.E.EM.ES.TDL.T.D.SRNG.....PV.K.NV.IP.N.IYSD.YEKG.LTNS.KK.MV.M.....N.D.GD.K...LS...Y...					536
4.11	EK.TA.N.E.EM.ES.TDL.T.D.SRNG.....PV.K.NV.IP.N.IYSD.YEKG.LTNS.KK.MV.M.....N.D.GD.K...LS...Y...					536
4.14	GK.TA.N.E.EM.ES.TDL.T.D.SRNG.....PV.K.NV.IP.N.IYSD.YEKG.LTNS.KK.MV.M...KN.D.GD.K...LS...Y...					536
5.3N.....D.....I.N.....G.....D.....Q.....					548
5.4N.....D.....I.N.....G.....D.....Q.....					548
5.6N.....D.....I.N.....G.....D.....Q.....					548
5.7N.....D.....I.N.....G.....R.....Q.....					548
5.8Q.N.....D.D.....I.N.....V.....M.....Q.....					548
6.1	DQ.EG.VTE...M.E.QI...KES.M.SNG...L.V...K...V.AP...I.D.V.KQ.TTHL.RV.V...K.E.ND.R.F.L...F...					547
6.2	DQ.EG.VTE...M.E.QI...KES.M.SNG...L.V...K...V.AP...I.D.V.KQ.TTHL.RV.V...K.E.ND.R.F.L...F...					547
6.4	DQ.EG.VTE...M.E.QI...KES.M.SNG...L.V...K...V.AP...I.D.V.KQ.TTHL.RV.V...K.E.ND.R.F.L...F...					547
6.5	DQ.EG.VTE...M.E.QI...KES.M.SNG...L.V...K...V.AP...I.D.V.KQ.TTHL.RV.V...K.E.ND.R.F.L...F...					547
6.6	DQ.EG.VTE...M.E.QI...KES.M.SNG...L.V...K...V.AP...I.D.V.KQ.TTHL.RV.V...K.E.ND.R.F.L...F...					547
6.7	DQ.EG.VTE...M.E.QI...KES.M.SNG...L.V...K...V.AP...I.D.V.KQ.TTHL.RV.V...K.E.ND.R.F.L...F...					547
7.5	.TVN...SDEERET.E.D.NI...KQ.KETKVN...S...TH...KT.EE.TP.MKL.HMEERR.IP.ST.L.VITIL.N.CSPLN.E...LSL...Y...T					544

	560	580	600	620	640	660																																									
Consensus	VLANRLKEVMPKI	IKTNQAYA	IKGRDIADVTMSI	KDTIDYMQEKKMNGFLI	SVDFEKA	FDRVEHTYLF	FDVLKTFGFGENFI	NWIKILYKGAFTKIKCNGFLTDCFKITRSI	659																																						
1.1	S	S	T	R	IND	QKD	L	DF	G	S	K	VQ	R	V	R	L	664																														
1.4	S	S	T	R	IND	QKD	L	DF	G	S	K	VQ	R	V	R	L	665																														
1.5	S	S	T	R	IND	QKD	L	DF	G	S	K	V	R	V	R	L	665																														
1.6	S	S	T	R	IND	QKD	L	DF	G	S	K	VQ	R	V	R	R	664																														
1.8	S	S	T	R	IND	QKD	L	DF	G	S	K	VQ	R	V	R	L	665																														
1.9	S	S	T	R	IND	QKD	L	DF	G	S	K	VQ	R	V	R	L	665																														
1.10	S	S	T	R	IND	QKD	L	DF	G	S	K	V	R	V	R	L	665																														
1.11	S	S	T	R	IND	QKD	L	DF	G	S	K	VQ	R	V	R	L	666																														
2.1	S	L	S	S	L	S	L	S	L	S	K	V	R	V	R	L	660																														
2.2	L	S	Y	S	L	S	L	S	L	S	N	V	E	V	E	L	660																														
2.3	L	S	Y	S	L	S	L	S	L	S	N	V	E	V	E	L	656																														
2.5	L	S	Y	S	L	S	L	S	L	S	N	V	E	V	E	L	660																														
2.7	L	S	Y	S	L	S	L	S	L	S	N	V	E	V	E	L	660																														
2.8	L	S	Y	S	L	S	L	S	L	S	N	V	E	V	E	L	660																														
2.10	L	S	Y	S	L	S	L	S	L	S	N	V	E	V	E	L	660																														
2.11	L	S	Y	S	L	S	L	S	L	S	N	V	E	V	E	L	660																														
S2.1	L	S	Y	S	L	S	L	S	L	S	N	V	E	V	E	L	660																														
S2.2	L	S	Y	S	L	S	L	S	L	S	N	V	E	V	E	L	660																														
3.1	I	T	I	I	E	K	EE	YI	L	QF	K	K	V	RK	IL	E	656																														
3.3	I	T	I	I	E	K	EE	YI	L	QF	K	K	V	RK	IL	E	656																														
3.4	I	T	I	I	E	K	EE	YI	L	QF	K	K	V	RK	IL	E	656																														
4.5	V	V	N	VLK	T	INN	R	I	W	R	E	ET	YI	L	K	MM	MER	RYLQ	C	ADITSC	V	KD	647																								
4.10	V	V	N	VLK	T	INN	R	I	W	R	E	ET	YI	L	K	MM	MER	RYLQ	C	ADITSC	V	KD	647																								
4.11	V	V	N	VLK	T	INN	R	I	W	R	E	ET	YI	L	K	MM	MER	RYLQ	C	ADITSC	V	KD	647																								
4.14	V	V	N	VLK	T	INN	R	I	W	R	E	ET	YI	L	K	MM	MER	RYLQ	C	ADITSC	V	KD	647																								
5.3	L	S	R	S	L	S	L	S	L	S	L	V	L	V	L	V	L	V	L	V	L	V	659																								
5.4	L	S	R	S	L	S	L	S	L	S	L	V	L	V	L	V	L	V	L	V	L	V	659																								
5.6	L	S	R	S	L	S	L	S	L	S	L	V	L	V	L	V	L	V	L	V	L	V	659																								
5.7	L	S	R	S	L	S	L	S	L	S	L	V	L	V	L	V	L	V	L	V	L	V	659																								
5.8	L	S	R	S	L	S	L	S	L	S	L	V	L	V	L	V	L	V	L	V	L	V	659																								
6.1	K	L	D	S	GV	K	T	IIS	R	V	Q	K	RK	YV	T	NF	E	NKV	N	LR	C	N	ISR	V	I	NV	658																				
6.2	K	L	D	S	GV	K	T	IIS	R	V	Q	K	RK	YV	T	NF	E	NKV	N	LR	C	N	ISR	V	I	NV	658																				
6.4	K	L	D	S	GV	K	T	IIS	R	V	Q	K	RK	YV	T	NF	E	NKV	N	LR	C	N	ISR	V	I	NV	658																				
6.5	K	L	D	S	GV	K	T	IIS	R	V	Q	K	RK	YV	T	NF	E	NKV	N	LR	C	N	ISR	V	I	NV	658																				
6.6	K	L	D	S	GV	K	T	IIS	R	V	Q	K	RK	YV	T	NF	E	NKV	N	LR	C	N	ISR	V	I	NV	658																				
6.7	K	L	D	S	GV	K	T	IIS	R	V	Q	K	RK	YV	T	NF	E	NKV	N	LR	C	N	ISR	V	I	NV	658																				
7.5	I	I	R	VRS	SP	HG	P	T	T	I	C	T	R	VVSS	DKD	GEG	L	M	L	C	I	L	N	NF	ME	Q	A	M	R	R	Y	K	I	L	N	K	S	C	V	V	N	T	P	L	E	V	655

	680	700	720	740	760	
Consensus	RQGCPLSALLYSLVAEPLGLA	IKQETKIKG	IKIEEEDEGKIYQYADDTTL	IVKDKKSVKEAMKKVQEFCKGTGAK	INENKTVYMRFGKADIL	TDCFQFKEVE-EIKILGI 769
1.1	A	V.H.DR	EV-GGVN-F	LQ.LA	Q.EI.H	S.D.L.RTEA.SGH.T.MD- 772
1.4	A	V.H.DR	EV-GGVN-F	LQ.LA	Q.ETI.H	S.D.L.RTEA.SGH.T.MD- 773
1.5	A	V.H.DI	EV-GGVN-F	L.LA	K.ET.H	S.D.L.ETEA.SGH.T.MD- 773
1.6	A.K	V.H.DR	EV-GGVN-F	LQELA	K.ET.H	S.D.G.L.RTEA.SGH.T.MD- 772
1.8	A	V.H.DR	EV-GGVN-F	LQ.LA	Q.ETI.H	S.D.L.RTEA.SGH.T.MD- 773
1.9	A	V.H.DR	EV-GGVN-F	LQ.LA	Q.ETI.H	S.D.L.RTEA.SGH.T.MD- 773
1.10	A	V.H.DI	EV-GGVN-F	L.LA	K.ET.H	S.D.L.ETEA.SGH.T.MD- 773
1.11	A	V.H.DR	EV-GGVN-F	LQ.LA	Q.ETI.H	S.D.L.RTEA.SGH.T.MD- 774
2.1	I			E		S.Q.R.L 770
2.2	I			E		S.Q.R.L 770
2.3	I			E		S.Q.R.L.M 766
2.5				E		S.Q.R.L 770
2.7	I	T		E		S.Q.R.R.L 770
2.8	I	T		E		S.Q.R.R.L 770
2.10	I			E		S.Q.R.L 770
2.11	I			E		S.Q.R.L 770
S2.1	I			E		S.Q.R.L 770
S2.2	I			E		S.Q.R.L 770
3.1		NE.E	E.NKVNK.MF	G.E	I.Q	S.S.V.D.K.ETD.AE.TN.V 766
3.3		NE.E	E.NKVNK.MF	G.E	K.I.Q	S.S.V.D.K.ETD.AE.TN.V 766
3.4		NE.E	E.NKVNK.MF	G.E	I.Q	S.S.V.D.K.ETD.AE.TN.V 766
4.5	M	T.S.T	DN.Q.R.LC.KGTDL.H.F	I	IEK.EEILNRY.N	KE.K.L.INV.PEKIK.ETVDM. 758
4.10	M	T.S.T	DN.Q.R.LC.KGTDL.H.F	I	IEK.EEILNRY.N	KE.K.L.INV.PEKIK.EKVDM. 758
4.11	M	T.S.T	DN.Q.R.LC.KGTDL.H.F	I	IEK.EEILNRY.N	KE.K.L.INV.PEKIK.EKVDM. 758
4.14	M	T.S.T	DN.Q.R.LC.KGTDL.H.F	I	IEK.EEILNRY	KE.K.L.INV.PEKIK.EKVDM. 758
5.3	I			E		S.Q.R.L 769
5.4	I			E		S.Q.R.R 769
5.6	I			E		S.Q.R.L 769
5.7	I			E		S.Q.R.L 769
5.8	I		D	E		S.Q.R.L 769
6.1	K	Q.T.A.S	IRDKE.H.V.T.E.Q	L.MQ	IN.IIERY.N.S	V.ME.V.L.N.C.P.V.K.DQK- 768
6.2	K	Q.T.A.S	IRDKE.H.V.T.E.Q	L.MQ	IN.IIERY.N.S	V.ME.V.L.N.C.P.V.K.DQK- 768
6.4	K	Q.T.A.S	IRDKE.H.V.T.E.Q	L.IQ	IN.IIERY.N.S	V.ME.V.L.N.C.P.V.K.DQK- 768
6.5	K	Q.T.A.S	IRDKE.H.V.T.E.Q	L.IQ	IN.IIERY.N.S	V.ME.V.L.N.C.P.V.K.DQK- 768
6.6	K	Q.T.A.S	IRDKE.H.V.T.E.Q	L.MQ	IN.IIERY.N.S	V.ME.V.L.N.C.P.V.K.DQK- 768
6.7	K	Q.T.A.S	IRDKE.H.V.T.E.Q	L.MQ	IN.IIERY.N.S	V.ME.V.L.N.C.P.V.K.DQK- 768
7.5	M	ISI.S.ATLV	.DKE.R.Q.PYGGVS-L.H	FT.R.MR.I	RI.NHIEVYG.AS	.IE.SEI.SI.GVGLSGCDIP.VAKEQTV.V 765

	900	920	940	960	980	
Consensus	K I V K K Y L Q E E N K A E W K K V M K Y F L N K V G N F N L G E D I L W M K T K N W M T E K L P G F Y Q E I L S A W G K F L D G V F Y Q V K G R E N L L N Q P L F L N K S I L K Q G K E L F F K K W M E V G I L R I R D V L					991
1.1	.M.R..D.D..A..RT.E..S.S.....DN..C.RM.KF..G..D..K.LIG.....TC.HFN IQ...I.....SG..N.E.VV..R..W...T.V....					994
1.4	.M.R..D.D..A..RT.E..S.S.....M.DN..C.RM.KF..G..D..K.LIG.....TC.HFN IQ...I.....SG..N.E.VV..R..W...T.V....					995
1.5	.M.R..D.D..A..RT.E..S.S.....DN..C.RM.KF..G..D..K.LIG.....TC.HFN IQ...I.....SG..N.E.VV..R..W...T.V....					995
1.6	.M.R..D.D..A..RT.E..S.S.....DN..Y.RM.KF..G..D..K.LIG.....TC.HFN IQ...I.....SG..N.E.VV..R..W...T.V....					994
1.8	.M.R..D.D..A..RT.E..S.S.....M.DN..C.RM.KF..G..D..K.LIG.....TC.HFN IQ...I.....SG..N.E.VV..R..W...T.V....					995
1.9	.M.R..D.D..A..RT.E..S.S.....M.DN..C.RM.KF..G..D..K.LIG.....TC.HFN IQ...I.....SG..N.E.VV..R..W...T.V....					995
1.10	.M.R..D.D..A..RT.E..S.S.....DN..C.RM.KF..G..D..K.LIG.....TC.HFN IQ...I.....SG..N.E.VV..R..W...T.V....					995
1.11	.M.R..D.D..A..RT.E..S.S.....DN..C.RM.KF..G..D..K.LIG.....TC.HFN IQ...I.....SG..N.E.VV..R..W...T.V....					996
2.1L.....ST.....L.....E.....D.....					992
2.2I.....L.....ST.....L.....E.....D.....					992
2.3L.....L.....ST.....L.....E.....D.....					988
2.5L.....L.....ST.....L.....E.....D.....					992
2.7I.....L.....L.....ST.....L.....E.....D.....					992
2.8I.....L.....L.....ST.....L.....E.....D.....					992
2.10L.....L.....L.....ST.....L.....E.....D.....					992
2.11L.....L.....L.....ST.....L.....E.....D.....					992
S2.1L.....L.....L.....ST.....L.....E.....D.....					992
S2.2L.....L.....L.....ST.....L.....E.....D.....					992
3.1	.R.....E.K.....T.....C.....M.DG.....A.....N.E.R.....G.....N.KQ.E.SPH...I.....NN.S...V.YY...I...KV..I					988
3.3	.R.I.....E.K.....T.....C.....M.DG.....A.....N.E.R.....N.KQ.E.SPH...I.....NN.S...V.YY...I...KV..I					988
3.4	.R.I.....E.K.....T.....C.....M.DG.....A.....N.E.R.....N.KQ.E.SPH...I.....NN.S...V.YY...I...KV..I					988
4.5	.T.N..WK..-..HV..D..R.....C--EIQ.Y.....P.EN.MN.V.E..K.VIK..D.RKHIDCSF-SK.EI.K.....EY.KEGNNTI.Y...FNA..KQ.K.I					976
4.10	.T.N..WK..-..HV..D..R.....C--EMQ.Y.....P.EN.MN.V.E..K.VIK..D.RKHIDCSF-SK.EI.K.....EY.KEGNNTI.Y...FNA..KQ.K.I					976
4.11	.T.N..WK..-..HV..D..R.....C--EMQ.Y.....P.EN.MN.V.E..K.VIK..D.RKHIDCSF-SK.EI.K.....EY.KEGNNTI.Y...FNA..KQ.K.I					976
4.14	.T.N..WK..-..HV..D..R.....C--EMQ.Y.....P.EN.MN.V.E..K.VIK..D.RKHIDCSF-SK.EI.K.....EY.KEGNNTI.Y...FNA..KQ.K.I					976
5.3L.....L.....L.....ST.....L.....E.....D.....					991
5.4L.....L.....L.....ST.....L.....E.....D.....					991
5.6L.....L.....L.....ST.....L.....E.....D.....					991
5.7L.....L.....L.....ST.....L.....E.....D.....					991
5.8L.....L.....L.....ST.....L.....E.....D.....					991
6.1	.V.I.....NKKF.....H..RFY..C.G..MND.....L.QS.ITGI.E..R.V.Q..SE..NH.DFKPE..NM.....AN..T.E..VY..Y.L.A.VCQVK...990					990
6.2	.V.I.....NK.F.....H..RFY..C.G..MND.....L.QS.ITGI.E..R.V.Q..SE..NH.DFKPE..NM.....GN..T.E..VY..Y.L.A.VCQVK...990					990
6.4	.V.I.....NK.F.....H..RFY..C.G..MND.....L.QS.ITGI.E..R.V.Q..SE..NH.DFKPE..NM.....GN..T.E..IY..Y.L.A.VCQVK...990					990
6.5	.V.I.....NK.F.....H..RFY..C.G..MND.....L.QS.ITGI.E..R.V.Q..SE..NH.DFKPE..NM.....GN..T.E..VY..Y.L.A.VCQVK...990					990
6.6	.V.I.....NK.F.....H..RFY..C.G..MND.....L.QS.ITGI.E..R.V.Q..SE..NH.DFKPE..NM.....GN..T.K..VY..Y.L.A.VCQVK...990					990
6.7	.V.I.....NK.F.....H..RFY..C.G..MND.....L.QS.ITGI.E..R.V.Q..SE..NH.DFKPE..NM.....GN..T.K..VY..Y.L.A.VCQVK...990					990
7.5	.M..L.YGNEEFG..GF.REY.S.SSG--C..EGVF.SL.KP.Y.NI.V...VC...SE.VME.G.ECENINQIHK..V...PK.QMR.RMFYNRLY.KA..RQVK.MV					985

	1,000	1,020	1,040	1,060	1,080	1,100	
Consensus	YEFKEGFLPKQYIVDL	MEEAKEEYSRKEI	EKKLEIVKGAIPKEWITRIENME-	EXGNEKVVHYYL-K-	EKLCNFKDCLLKDFYVYFRD	SVFKEPIANNFWVQRXNSVKKEN	1099
1.1QV..VI.V.D...	DFN.QDLI.EYD.I.N...	A..L.....NKQS.D.I.KF-G-	WW.L..STV.MI.GF..G...	K.RS.EY.IRMFKD.NED.		1102
1.4V..VI.V.D...	DFN.QDLI.EYD.I.N...	A..L.....NKQS.D.I.KF-G-	WW.L..STV.MI.GF..G...	K.RS.EY.IRMFKD.NED.		1103
1.5V..VI.V.D...	DFN.QDLI.EYD.I.N...	A..L.....NKQS.D.I.RF-G-	WW.L..STV.MI.GF..G...	K.RS.EY.IRMFKD.NED.		1103
1.6V..VI.V.D...	DFN.QDLI.EYD.I.N...	A..L.....NKHS.DMI.RF-G-	YW.L..STV.MI.GF..G...	K.RS.EY.IRMFKD.NED.		1102
1.8V..VI.V.D...	DFN.QDLI.EYD.I.N...	A..L.....NKQS.D.I.KF-G-	WW.L..STV.MI.GF..G...	K.RS.EY.IRMFKD.NED.		1103
1.9V..VI.V.D...	DFN.QDLI.EYD.I.N...	A..L.....NKQS.D.I.KF-G-	WW.L..STV.MI.GF..G...	K.RS.EY.IRMFKD.NED.		1103
1.10V..VI.V.D...	DFN.QDLI.EYD.I.N...	A..L.....NKQS.D.I.RF-G-	WW.L..STV.MI.GF..G...	K.RS.EY.IRMFKD.NED.		1103
1.11V..VI.V.D...	DFN.QDLI.EYD.I.N...	A..L.....NKQS.D.I.RF-G-	WW.L..STV.MI.GF..G...	K.RS.EY.IRMFKD.NED.		1104
2.1	..K...T.....V...N...T.....C.....I.....G.....D.....Q.....L.....		1100
2.2	..K...T.....V...N...T.....G.....I...N...G.....D.....Q.....L.....		1100
2.3	..K...T.....V...N...T.....G.....I.....G.....D.....Q.....L.....		1096
2.5	..K...T.....V...N...T.....C.....I.....G.....D.....Q.....L.....		1100
2.7	..K...T.....V...N...T...S...G.....I.....G.....I.....Q.....L...N...		1100
2.8	..K...T.....V...N...T...S...G.....I.....G.....I.....Q.....L...N...		1100
2.10	..K...T.....V...N...T.....C.....I...C...G.....D.....Q.....L.....		1100
2.11	..K...T.....V...N...T.....C.....I.....G.....D.....Q.....L.....		1100
S2.1	..K...T.....V...N...T.....C.....I.....G.....D.....Q.....L.....		1100
S2.2	..K...T.....V...N...T.....C.....I.....G.....D.....Q.....L.....		1100
3.1TE..VI.T.....N.....N.D.I.Q.....	RS..KF--KEKET.E.Y.KT-G-	..I...NE.TV.NI.CF..EN...	T..KY.IEKYKN..ANE		1096
3.3TE..VI.T.....N.....N.D.I.Q.....	RS..F--KEKET.E.Y.KT-G-	..IG..NE.TV.NI.CF..EN...	T..KY.IEKYKN..ANE		1096
3.4TE..VI.T.....N.....N.D.I.Q.....	KS..F--KEKET.E.Y.KT-G-	..I...NE.TV.NI.CF..EN...	T..KY.IEKYKN..ANE		1096
4.5	..VIP.YVQA.V.K.A.II.NY.NE.KRV.	NQFNLLKN...KEI...KL-	MHRDEGRGINF--NQYA...	GI..M..SCL.KD.VT.K.EEY.	KRLYPNIE.KE		1084
4.10	..VIP.YVQAHV.K.A.II.NY.NETKRV.	Q.NNL.KN...KEI...KF-	MHSDEGRVINF--NQYA...	GI..M..SCL.KD.VT.K.EEY.	KRLYPNIE.KE		1084
4.11	..VIP.YVQAHV.K.A.II.NY.NETKRV.	Q.NNL.KN...KEI...KF-	MHSDEGRVINF--NQYA...	GI..M..SCL.KD.VT.K.EEY.	KRLYPNIE.KE		1084
4.14	..VIP.YVQAHV.K.A.II.NY.NETKRV.	Q.NNL.KN...KEI...KL-	MHRDEGRVINF--NQYA...	GI..M..SCLCKD.VT.K.EEY.	KRLYPNIE.KE		1084
5.3	..K...N.....V...N...T.....G.....I.....G.....D.....Q.....L...E...		1099
5.4	..K...N.....V...N...T...S...G.....I...N...G.....D.....Q.....L...E...		1099
5.6	..K...N.....V...N...T.....G.....I.....G.....D.....Q.....L...E...		1099
5.7	..K...T.....V...N...T.....G.....I...N...G.....D.....Q.....L...E...		1099
5.8	..K...T.....V...N...T.....C..G..I.....G.....D.....Q.....L...E...		1099
6.1	..I.....S.V...A.DGIGKDIK.ETLV.QF...	S.L...VK.QNN-S-DN-K.F--	EKD.QE.V..N..IC..NK.H.R.VSEQ..	KKIFVEIDE.G			1096
6.2	..I.....S.V...A.DGIGKDIK.ETLV.QF...	S.L...VK.QNN-S-DN-K.F--	EKD.QE.V..N..IC..NK.H.R.VSEQ..	KKIFVEIDE.G			1096
6.4	..I.....S.V...A.DGIGKDIK.ETLV.QF...	S.L...VK.QNN-S-DN-K.F--	EKD.QE.V..N..IC..NK.H.R.VSEQ..	KKIFVEIDE.G			1096
6.5	..I.....S.V...A.DGIGKDIK.ETLV.QF...	S.L...VK.QNN-S-DN-K.F--	EKD.QE.V..N..IC..NK.H.R.VSEQ..	KKIFVEIDE.G			1096
6.6	..I.....S.V...A.DGIGKDIK.ETLV.QF...	S.L...VK.QNN-S-DN-K.F--	EKD.QE.V..N..IC..NK.H.R.VSEQ..	KKIFVEIDE.G			1096
6.7	..I.....S.V...A.DGIGKDIK.ETLV.QF...	S.L...VK.QNN-S-DN-K.F--	EKD.QE.V..N..IC..NK.H.R.VSEQ..	KKIFVEIDE.G			1096
7.5	..VK...NRA.Y.SVV.WDD.IKKSQVDSVC.NIRVSL..R.VEM..RESARP..WGMPEMFIV.NG.KICLS.VIT.TI.GI.VEKEI.L.ASE.V.PKLFPNLNVKT						1096

	1,120	1,140	1,160	1,180	1,200	1,220		
Consensus	IWKNM	RGKII	ETKLECFEYFIRHKAIFTEC	ILTKIGMEENATCKVCFQEDEGI	LHLFLYCKE	LECFYXKCKKMLKDLL-	KXWDEEELEWXTLVMFGWNMENKKNKFFVNLLI 1209	
1.1	A.IK.LVQS.V.NL.L.N.V.DI.N	SV	QDA.F	N.KD.NE.Y.SIILT.K-GER.D	EKVL.L.V.K.CN.E.LI	V	1211	
1.4	A.IN.LVQS.V.NL.L.N.V.DI.N	SV	QDA.F	N.KD.NE.SIILT.K-GER.D	EKVL.L.V.K.CN.E.LI	V	1212	
1.5	A.IN.LVQS.V.NL.L.N.V.DI.N	SV	QDA.F	N.KD.NE.SIILT.K-GER.D	EKVL.L.V.K.CN.E.LI	V	1212	
1.6	A.IK.LVQS.V.NL.L.N.V.DI.N	SV	QDA.F	N.KD.NE.SIILT.K-GER.D	EKVL.L.V.K.CN.E.LI	V	1211	
1.8	A.IN.LVQS.V.NL.L.N.V.DI.N	SV	QDA.F	N.KD.NE.SIILT.K-GER.D	EKVL.L.V.K.CN.E.LI	V	1212	
1.9	A.IN.LVQS.V.NL.L.N.V.DI.N	SV	QDA.F	N.KD.NE.SIILT.K-GER.D	EKVL.L.V.K.CN.E.LI	V	1212	
1.10	A.IN.LVQS.V.NL.L.N.V.DI.N	SV	QDA.F	N.KD.NE.SIILT.K-GER.D	EKVL.L.V.K.CN.E.LI	V	1212	
1.11	A.IK.LEQS.V.NL.L.N.V.DI.N	SV	QDA.F	N.KD.NE.SIILT.K-GER.D	EKVL.L.V.K.CN.E.LI	V	1213	
2.1	R	HI.Q	K.Q	D	Q	N	1210	
2.2	R	QR.P	M.Q	D	Q	N	1210	
2.3	I	QR.P	K.Q	D	Q	N	1206	
2.5	R	HI.Q	M.Q	D	Q	N	1210	
2.7	I	QR.H	M.Q	D	DQ	N	1210	
2.8	I	QR.H	M.Q	D	DQ	N	1210	
2.10	R	HI.Q	K.Q	D	Q	N	1210	
2.11	R	HI.Q	K.Q	D	Q	N	1210	
S2.1	R	HI.Q	K.Q	D	Q	N	1210	
S2.2	R	HI.Q	K.Q	D	Q	N	1210	
3.1	G.K.RYV.V.L.FL	SDV.N	QSGM.QK.E.F.M	QK.G.LKD.VLI.G	GD.N.T.NRV.L	KN.E.I.C	1206	
3.3	G.K.RYV.V.L.FL	SDV.N	QSGM.QK.E.F.M	Q.G.LKD.VLI.G	GD.N.T.NRV.L	KN.E.I.C	1206	
3.4	G.K.RYV.V.L.FL	SDV.N	QSGM.QK.E.F.M	Q.G.LKD.VLI.G	GD.N.T.NRV.L	KN.E.I.C	1206	
4.5	Q.L.AWWKSPV.N.D.I.L.QNCLLS.MR.C	LAND.R	NE	FK.K.IG.I.E.NEMG-IENEVVDE	E.FL.CKT.G	TR.I.FIL	1194	
4.10	QHV.AWWKSPV.N.D.I.L.QNCLLS.MR.C	LAND.R	NE	FK.K.IG.I.E.NEMG-IENEVVDE	E.FL.CKT.G	TR.I.FFL	1194	
4.11	QHV.AWWKSPV.N.D.I.L.QNCLLS.MR.C	LAND.R	NE	FK.K.IG.I.E.NEMG-IENEVVDE	E.FL.CKT.G	TR.I.FFL	1194	
4.14	Q.L.AWWKSPV.N.D.I.L.QNCLLS.MR.C	LAND.R	NE	FK.K.IG.I.E.NEMG-IENEVVDD	E.FL.CKT.G	TR.M.FIL	1194	
5.3	R	R	QR.P	M.Q	D	Q	N	1209
5.4	R	HI.Q	M.Q	D	Q	N	1209	
5.6	R	R	QR.P	M.Q	D	Q	N	1209
5.7	E	R	QR.P	M.Q	D	QF	N	1209
5.8	R	QR.P	D	M.Q	D	Q	N	1209
6.1	LKW.YLDA.I.MLD	NV.TK.AQ.SS	L.I.GL.N.M	K.V.NT.MA.L.E.M.QMMASETEQI	E.TEWYKI	LFGFNGSKTRIA	FL 1207	
6.2	LKW.YLDA.I.MLD	NV.TK.AQ.SS	L.I.GL.N.M	K.V.NT.MA.L.E.M.QMMPSETEQI	EDTEWYKI	LFGFNGSKTRIA	FL 1207	
6.4	LKW.YLDA.I.MLD	NV.TK.AQ.SS	L.I.GL.N.M	K.V.NT.MA.L.E.M.QMMPSETEQI	EDTEWYKI	LFGFNGSKTRIA	FL 1207	
6.5	LKW.YLDA.I.MLD	NV.TK.AQ.SS	L.I.GL.N.M	K.V.NT.MA.L.E.M.QMMPSETEQI	EDTEWYKI	LFGFNGSKTRIA	FL 1207	
6.6	LKW.YLDA.I.MLD	NV.TK.AQ.SS	L.I.GL.N.M	K.V.NT.MA.L.E.M.QMMPSETEQI	EDTEWYKI	LFGFNGSKTRIA	FL 1207	
6.7	LKW.YLDA.I.MLD	NV.TK.AQ.SS	L.I.GL.N.M	K.V.NT.MA.L.E.M.QMMPSETEQI	EDTEWYKI	LFGFNGSKTRIA	FL 1207	
7.5	LNV.YNSLDC.NLDFKL	NR.Y.KVVIHQ.NRNV	RE.D.KSDP.TLM.I.FE	GF.HE.L.QLI	NW-GKNAQG-N	KQ.FL.E.TNC.E.N	CN 1204	

	1,240	1,260	1,280	1,300																		
Consensus	MMI	KKC	IWERR	NAKQEKVV	LNWVNF	KRKMERY	I	EXL	YFYFK	GEDML	PSFYD	VFNDE	EVYN	VLNGL	-	KWKMPKE	-----	1282				
1.1	LM	SAV	VA	K	A	D		V	K	V	C	F	L	QEA	TQ	SKI	DT	GM	F	-----	1283	
1.4	LM	SA	VA	K	A	D		V	K	V	C	F	L	QEA	TQ	SKI	DT	GL	F	-----	1284	
1.5	LM	SA	VA	K	A	D		V	K	V	FVLLF									-----	1253	
1.6	LM	SA	VA	K	A	DA		V	K	V	C	F	L	QEA	TQ	SKI	DT	GM	F	-----	1283	
1.8	LM	SA	VA	K	A	D		V	K	V	C	F	L	QEA	TQ	SKI	DT	GL	F	-----	1284	
1.9	LM	SA	VA	K	A	D		V	K	V	C	F	L	QEA	TQ	SKI	DT	GL	F	-----	1284	
1.10	LM	SA	VA	K	A	D		V	K	V	FVLLF									-----	1253	
1.11	LM	SA	VA	K	A	FD		V	K	V	C	F	L	QEA	TQ	SKI	DT	GM	F	-----	1285	
2.1	V																			-----	1233	
2.2	V							I		L											-----	1287
2.3	V							L													-----	1283
2.5	V																				-----	1233
2.7	V							I		L											-----	1287
2.8	V							I		L											-----	1287
2.10	V							K													-----	1287
2.11	V							L													-----	1287
S2.1	V							L													-----	1287
S2.2	V							L													-----	1287
3.1	LM	NAM	I	K	M	MD	GI		V	K	V	H	N	NI	SELHKILT	KACQ	FKEM	N	L	HF	-----	1279
3.3	LM	NAM	I	K	M	MD	GI		V	K	V	H	N	NI	SELHKILT	KACQ	FKEM	N	L	HF	-----	1279
3.4	LM	NAM	I	K	M	MD	GI		V	K	I	H	N	NI	SELHKILT	KACQ	FKEM	N	L	HF	-----	1279
4.5	TVARNV	N	IV	KNGK	S	CKL	Q	SELLNT	F	CNMN	KI	DV	FEKDFL	SG	TYVQMTLY	NVNVLL	PECNCF				-----	1273
4.10	TVARNV	N	IV	KNGK	S	CKL	Q	SELLNT	F	CNMN	KI	DV	FEKDFL	NG	TYVQMALYN	VNVLL	PECNCF				-----	1273
4.11	TVARNV	N	IV	NNGK	S	CKL	Q	SELLNT	F	CNMN	KI	DV	FEKDFL	NG	TYVQMALYN	VNVLL	PECNCF				-----	1273
4.14	TVARNV	N	IV	KNGK	S	CKL	Q	SELLNT	F	CNMN	KI	DV	FEKDFL	NG	TYVQMALYN	VNVLL	PECNCF				-----	1273
5.3	V																				-----	1286
5.4	V																				-----	1242
5.6	V																				-----	1286
5.7	V																				-----	1286
5.8	V																				-----	1286
6.1	SVARTV		IV	NKNAK	IEL	KMY	K	L	N	KTVEQ	RM	EKETE	KT	MLNTPY	IMYT	DGL	IDLPE				-----	1282
6.2	SIARTV		IV	NKNAK	IEL	KMY	K	L	N	KTVEQ	RM	EKETE	KT	MLNTPY	IMYT	DGL	IDLPERLCLMF	S	CFNTN	IMS	-----	1297
6.4	SVARTV		IV	NKNAK	IEL	KMY	KNWKT	I	LKLWNN	I	LEWKKR	LN	TKRLC								-----	1263
6.5	SVARTV		IV	NKNAK	IEL	KMY	K	L	N	KTVEQ	RM	EKETE	KT	MLNTPY	IMYT	DGL	IDLPE				-----	1282
6.6	SVARTV		IV	NKNAK	IEL	KMY	K	L	N	KTVEQC	RM	EKETE	KT	MLNTPY	IMYT	DGL	IDLPE				-----	1282
6.7	SVARTV		IV	NKNAK	IEL	KMY	K	L	N	KTVEQC	RM	EKETE	KT	MLNTPY	IMYT	DGL	IDLPE				-----	1282
7.5	FFLSHARY	-	A	IW	RRNMAY	F	EAKI	IDVNV	FKSVVR	KNV	LMWKY	LRKDV	FEEAF	I	HE	E	ITV	DH	KGLMLNF		-----	1277